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Sequence Listing could not be accepted due to errors.

See attached Validation Report.

If you need help call the Patent Electronic Business Center at (866)
217-9197 (toll free).

Reviewer: Keisha Douglas

Timestamp: [year=2010; month=11; day=3; hr=10; min=53; sec=1; ms=998;]

=====

Reviewer Comments:

<210> 1

<211> 7

<212> PRT

<213> Gymnea sylvestre

<400> 1

Asn Gly Ser Phe Ser Gly Phe

1

5

The above sequence id# 1 is invalid, please delete spacing between the
amino acids and the numbering. This error is seen globally throughout
the sequences.

Application No: 10579655 Version No: 4.0

Input Set:**Output Set:**

Started: 2010-10-25 19:40:21.780
Finished: 2010-10-25 19:40:36.460
Elapsed: 0 hr(s) 0 min(s) 14 sec(s) 680 ms
Total Warnings: 379
Total Errors: 2984
No. of SeqIDs Defined: 415
Actual SeqID Count: 415

Error code	Error Description
W 402	Undefined organism found in <213> in SEQ ID (1)
E 355	Empty lines found between the amino acid numbering and the
E 321	No. of Bases conflict, this line has no nucleotides SEQID (1)
W 402	Undefined organism found in <213> in SEQ ID (2)
E 355	Empty lines found between the amino acid numbering and the
E 321	No. of Bases conflict, this line has no nucleotides SEQID (2)
W 402	Undefined organism found in <213> in SEQ ID (3)
E 355	Empty lines found between the amino acid numbering and the
E 321	No. of Bases conflict, this line has no nucleotides SEQID (3)
W 402	Undefined organism found in <213> in SEQ ID (4)
E 355	Empty lines found between the amino acid numbering and the
E 321	No. of Bases conflict, this line has no nucleotides SEQID (4)
W 213	Artificial or Unknown found in <213> in SEQ ID (5)
E 355	Empty lines found between the amino acid numbering and the
E 321	No. of Bases conflict, this line has no nucleotides SEQID (5)
E 355	Empty lines found between the amino acid numbering and the
E 321	No. of Bases conflict, this line has no nucleotides SEQID (5)
W 213	Artificial or Unknown found in <213> in SEQ ID (6)
E 355	Empty lines found between the amino acid numbering and the
E 321	No. of Bases conflict, this line has no nucleotides SEQID (6)

Input Set:

Output Set:

Started: 2010-10-25 19:40:21.780
Finished: 2010-10-25 19:40:36.460
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Total Warnings: 379
Total Errors: 2984
No. of SeqIDs Defined: 415
Actual SeqID Count: 415

Error code	Error Description
E 355	Empty lines found between the amino acid numbering and the
E 321	No. of Bases conflict, this line has no nucleotides SEQID (6)
W 213	Artificial or Unknown found in <213> in SEQ ID (7)
E 355	Empty lines found between the amino acid numbering and the
E 321	No. of Bases conflict, this line has no nucleotides SEQID (7)
E 355	Empty lines found between the amino acid numbering and the
E 321	No. of Bases conflict, this line has no nucleotides SEQID (7)
W 213	Artificial or Unknown found in <213> in SEQ ID (8)
E 355	Empty lines found between the amino acid numbering and the
E 321	No. of Bases conflict, this line has no nucleotides SEQID (8)
E 355	Empty lines found between the amino acid numbering and the
E 321	No. of Bases conflict, this line has no nucleotides SEQID (8)
W 402	Undefined organism found in <213> in SEQ ID (9)
E 355	Empty lines found between the amino acid numbering and the
E 321	No. of Bases conflict, this line has no nucleotides SEQID (9)
E 355	Empty lines found between the amino acid numbering and the
E 321	No. of Bases conflict, this line has no nucleotides SEQID (9)
W 402	Undefined organism found in <213> in SEQ ID (10)
E 355	Empty lines found between the amino acid numbering and the
E 321	No. of Bases conflict, this line has no nucleotides SEQID (10)
E 355	Empty lines found between the amino acid numbering and the
E 321	No. of Bases conflict, this line has no nucleotides SEQID (10)

Input Set:

Output Set:

Started: 2010-10-25 19:40:21.780
Finished: 2010-10-25 19:40:36.460
Elapsed: 0 hr(s) 0 min(s) 14 sec(s) 680 ms
Total Warnings: 379
Total Errors: 2984
No. of SeqIDs Defined: 415
Actual SeqID Count: 415

Error code	Error Description
W 402	Undefined organism found in <213> in SEQ ID (11)
E 355	Empty lines found between the amino acid numbering and the
E 321	No. of Bases conflict, this line has no nucleotides SEQID (11)
E 355	Empty lines found between the amino acid numbering and the
E 321	No. of Bases conflict, this line has no nucleotides SEQID (11)
E 355	Empty lines found between the amino acid numbering and the
E 321	No. of Bases conflict, this line has no nucleotides SEQID (11)
W 402	Undefined organism found in <213> in SEQ ID (12)
E 355	Empty lines found between the amino acid numbering and the proteins
E 321	No. of Bases conflict, this line has no nucleotides SEQID (12) POS (0)
W 402	Undefined organism found in <213> in SEQ ID (13)
W 402	Undefined organism found in <213> in SEQ ID (14)
W 402	Undefined organism found in <213> in SEQ ID (15)
W 402	Undefined organism found in <213> in SEQ ID (16)
W 402	Undefined organism found in <213> in SEQ ID (17)
W 213	Artificial or Unknown found in <213> in SEQ ID (18)
W 213	Artificial or Unknown found in <213> in SEQ ID (19)
W 213	Artificial or Unknown found in <213> in SEQ ID (20)
W 213	Artificial or Unknown found in <213> in SEQ ID (21)
W 213	Artificial or Unknown found in <213> in SEQ ID (22)
W 402	Undefined organism found in <213> in SEQ ID (23)

Input Set:

Output Set:

Started: 2010-10-25 19:40:21.780
Finished: 2010-10-25 19:40:36.460
Elapsed: 0 hr(s) 0 min(s) 14 sec(s) 680 ms
Total Warnings: 379
Total Errors: 2984
No. of SeqIDs Defined: 415
Actual SeqID Count: 415

Error code	Error Description
W 402	Undefined organism found in <213> in SEQ ID (24)
W 402	Undefined organism found in <213> in SEQ ID (25)
W 402	Undefined organism found in <213> in SEQ ID (26)
W 402	Undefined organism found in <213> in SEQ ID (28)
W 402	Undefined organism found in <213> in SEQ ID (29)
W 402	Undefined organism found in <213> in SEQ ID (30) This error has occurred more than 20 times, will not be displayed
W 213	Artificial or Unknown found in <213> in SEQ ID (77)
W 213	Artificial or Unknown found in <213> in SEQ ID (78)
W 213	Artificial or Unknown found in <213> in SEQ ID (79)
W 213	Artificial or Unknown found in <213> in SEQ ID (80)
W 213	Artificial or Unknown found in <213> in SEQ ID (253)
W 213	Artificial or Unknown found in <213> in SEQ ID (254)
W 213	Artificial or Unknown found in <213> in SEQ ID (255)
W 213	Artificial or Unknown found in <213> in SEQ ID (256)
W 213	Artificial or Unknown found in <213> in SEQ ID (257)
W 213	Artificial or Unknown found in <213> in SEQ ID (258)
W 213	Artificial or Unknown found in <213> in SEQ ID (259) This error has occurred more than 20 times, will not be displayed

SEQUENCE LISTING

<110> Sanofi Pasteur, Inc.

<120> METHODS FOR PURIFYING PERTUSSIS TOXIN AND PEPTIDES USEFUL
THEREFOR

<130> API-03-15

<140> 10579655

<141> 2010-10-25

<150> 60/523,881

<151> 2003-11-20

<150> PCT/US2004/038700

<151> 2004-11-18

<160> 415

<170> PatentIn version 3.5

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<211> 7

<212> PRT

<213> Gymnea sylvestre

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<210> 2

<211> 7

<212> PRT

<213> Gymnea sylvestre

<400> 2

Asn Gly Ser Phe Ser Gly Cys

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<210> 3

<211> 7

<212> PRT

<213> Gymnea sylvestre

<400> 3

Asp Gly Ser Phe Ser Gly Phe

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<210> 4

<211> 7

<212> PRT

<213> Gymnea sylvestre

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<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic sequence, no source organism

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Asn Met Arg Ile Glu Thr Pro Asn Asn Ile Arg Lys Asp Ala

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<223> Synthetic sequence, no source organism

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<210> 7

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<212> PRT

<213> Artificial Sequence

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Arg Ser Asn Val Ile Pro Leu Asn Glu Val Trp Tyr Asp Thr Gly Trp

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<210> 8

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<212> PRT

<213> Artificial Sequence

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<210> 9

<211> 34

<212> PRT

<213> Gymnea sylvestre

<400> 9

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Cys Cys Glu Pro Leu Glu Cys Ile Tyr Thr Ser Glu Leu Tyr Ala Thr

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Cys Gly

<210> 10

<211> 34

<212> PRT

<213> Gymnea sylvestre

<400> 10

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Cys Cys Glu Pro Leu Glu Cys Phe Gln Met Gly His Gly Phe Lys Arg

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Cys Gly

<210> 11

<211> 35

<212> PRT

<213> Gymnea sylvestre

<400> 11

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Cys Gly Ser

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<210> 12

<211> 34

<212> PRT

<213> Gymnea sylvestre

<400> 12

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20 25 30

Cys Gly

<210> 13

<211> 35

<212> PRT

<213> Gymnea sylvestre

<400> 13

Gln Gln Cys Val Lys Lys Asp Glu Leu Cys Ile Pro Tyr Tyr Leu Asp

1 5 10 15

Cys Cys Glu Pro Leu Glu Cys Lys Lys Val Asn Trp Trp Asp His Lys

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Cys Ile Gly

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<213> Gymnea sylvestre

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<211> 141

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<213> *Gymnea sylvestre*

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<221> misc_feature

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nnsnnstgct gtgagccct cgagtgcnn snnnsnnsnn nsnnnsnnsnn snnstgcggc 120

agcggcagtt ctgggtctag c 141

<210> 16

<211> 84

<212> DNA

<213> *Gymnea sylvestre*

<400> 16

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catagtggct caagctcagg atca

84

<210> 17

<211> 44

<212> DNA

<213> Gymnea sylvestre

<400> 17

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44

<210> 18

<211> 10

<212> RNA

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<223> Synthetic sequence, no source organism

<400> 18

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<210> 19

<211> 53

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20 25 30

Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Xaa Asp Ala Asn Ala Pro

35 40 45

Lys Ala Ser Ala Ile

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<210> 20

<211> 6

<212> PRT

<213> Artificial Sequence

<220>

<223> Synthetic sequence, no source organism

<400> 20

His His His His His His

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<210> 21

<211> 6

<212> PRT

<213> Artificial Sequence

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<210> 22

<211> 127

<212> DNA

<213> Artificial sequence

<220>

<223> Synthetic sequence, no source organism

<220>

<221> MISC_FEATURE

<222> (28)..(105)

<223> S represents C or G, and N represents A, T, G or C

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snnsnnsnns nnsnnsnnsn nsnnnsnnsn snnsnnsnns nnsnnagatc tagcatgatg 120

atgatga 127

<210> 23

<211> 81

<212> DNA

<213> Gymnea sylvestre

<400> 23

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catcatcatc atgctagatc t 81

<210> 24

<211> 32

<212> DNA

<213> *Gymnea sylvestre*

<400> 24

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<211> 18

<212> DNA

<213> Bacteriophage M13

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<400> 26

Met His His His His His Ser Gly Ser Ser Ser Gly Ser Gly Cys

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Val Lys Lys Asp Glu Leu Cys Ala Gly Ser Val Gly His Cys Cys Glu

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Pro Leu Glu Cys Leu Arg Arg Phe Leu Asn Leu Arg Trp Cys Gly Ser

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Gly Ser Ser Gly Ser Ser

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<210> 27

<211> 54

<212> PRT

<213> *Gymnema sylvestre*

<400> 27

Met His His His His His His Ser Gly Ser Ser Ser Gly Ser Gly Cys

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Val Lys Lys Asp Glu Leu Cys Ile Val Met Arg Ala Pro Cys Cys Glu

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Pro Leu Glu Cys Leu Arg Arg Tyr Met Leu Lys His Met Cys Gly Ser

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Gly Ser Ser Gly Ser Ser

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<210> 28

<211> 54

<212> PRT

<213> Gymnea sylvestre

<400> 28

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Val Lys Lys Asp Glu Leu Cys Lys Ala Phe Arg Tyr Ser Cys Cys Glu

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Pro Leu Glu Cys Leu Arg Lys Trp Leu Lys Ala Arg Phe Cys Gly Ser

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Gly Ser Ser Gly Ser Ser

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<210> 29

<211> 54

<212> PRT

<213> *Gymnea sylvestre*

<400> 29

Met His His His His His His Ser Gly Ser Ser Ser Gly Ser Gly Cys

1 5 10 15

Val Lys Lys Asp Glu Leu Cys Leu Arg Ser Ser Ile Asp Cys Cys Glu

20 25 30

Pro Leu Glu Cys Leu Tyr Lys Trp Met Gln Arg Arg Leu Cys Gly Ser

35 40 45

Gly Ser Ser Gly Ser Ser

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<210> 30

<211> 54

<212> PRT

<213> Gymnea sylvestre

<400> 30

Met His His His His His His Ser Gly Ser Ser Ser Gly Ser Gly Cys

1 5 10 15

Val Lys Lys Asp Glu Leu Cys Trp Pro Arg Arg His Lys Cys Cys Glu

20 25 30

Pro Leu Glu Cys Leu Leu Glu Met Leu Glu Arg Lys Arg Cys Gly Ser

35 40 45

Gly Ser Ser Gly Ser Ser

50

<210> 31

<211> 53

<212> PRT

<213> Gymnea sylvestre

<400> 31

Met His His His His His His Ser Gly Ser Ser Ser Gly Ser Gly Cys

1 5 10 15

Val Lys Lys Asp Glu Leu Cys Met Ser Met Ala Cys Val Cys Cys Glu

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25

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Pro Leu Glu Cys Lys Tyr His Gly Tyr Phe Trp Leu Cys Gly Ser Gly

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Ser Ser Gly Ser Ser

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<210> 32

<211> 54

<212> PRT

<213> *Gymnea sylvestre*

<400> 32

Met His His His His His His Ser Gly Ser Ser Ser Gly Ser Gly Cys

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Val Lys Lys Asp Glu Leu Cys Ala Val Trp Phe Asp Val Cys Cys Glu

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Pro Leu Glu Cys Thr Tyr Gln Ser Gly Tyr Tyr Trp Leu Cys Gly Ser

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Gly Ser Ser Gly Ser Ser

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<210> 33

<211> 54

<212> PRT

<213> Gymnea sylvestre

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Val Lys Lys Asp Glu Leu Cys Glu Pro Trp Tyr Trp Arg Cys Cys Glu

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Pro Leu Glu Cys Val Tyr Thr Ser Gly Tyr Tyr Tyr Ser Cys Gly Ser

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Gly Ser Ser Gly Ser Ser

<210> 34

<211> 54

<212> PRT

<213> *Gymnea sylvestre*

<400> 34

Met His His His His His His Ser Gly Ser Ser Ser Gly Ser Gly Cys

1 5 10 15

Val Lys Lys Asp Glu Leu Cys Ala Arg Trp Asp Leu Val Cys Cys Glu

20 25 30

Pro Leu Glu Cys Ile Tyr Thr Ser Glu Leu Tyr Ala Thr Cys Gly Ser

35 40 45

Gly Ser Ser Gly Ser Ser

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<210> 35

<211> 54

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<213> Gymnea sylvestre

<400> 35

Met His His His His His His Ser Gly Ser Ser Ser Gly Ser Gly Cys

1 5 10 15

Val Lys Lys Asp Glu Leu Cys Val Phe Tyr Phe Pro Asn Cys Cys Glu

20 25 30

Pro Leu Glu Cys Arg Trp Val Asn Asp Asn Tyr Gly Trp Cys Gly Ser

35 40 45

Gly Ser Ser Gly Ser Ser

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<210> 36

<211> 53

<212> PRT

<213> Gymnea sylvestre

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1 5 10 15

Val Lys Lys Asp Glu Leu Cys Met Ser Met Ala Cys Val Cys Cys Glu

20 25 30

Pro Leu Glu Cys Lys Tyr His Gly Tyr Phe Trp Leu Cys Gly Ser Gly

35 40 45

Ser Ser Gly Ser Ser

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<210> 37

<211> 54

<212> PRT

<213> Gymnea sylvestre

<400> 37

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1 5 10 15

Val Lys Lys Asp Glu Leu Cys Thr Thr Ala Ser Lys Ser Cys Cys Glu

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Pro Leu Glu Cys Lys Trp Thr Asn Glu His Phe Gly Thr Cys Gly Ser

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Gly Ser Ser Gly Ser Ser

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<210> 38

<211> 54

<212> PRT

<213> *Gymnea sylvestre*

<400> 38

Met His His His His His His Ser Gly Ser Ser Ser Gly Ser Gly Cys

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Val Lys Lys Asp Glu Leu Cys Ser Gln Ser Val Pro Met Cys Cys Glu

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Pro Leu Glu Cys Lys Trp Phe Asn Glu Asn Tyr Gly Ile Cys Gly Ser

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Gly Ser Ser Gly Ser Ser

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<210> 39

<211> 54

<212> PRT

<213> Gymnea sylvestre

<400> 39

Met His His His His His His Ser Gly Ser Ser Ser Gly Ser Gly Cys

1 5 10 15

Val Lys Lys Asp Glu Leu Cys Ala Arg Trp Asp Leu Val Cys Cys Glu

20 25 30

Pro Leu Glu Cys Ile Tyr Thr Ser Glu Leu Tyr Ala Thr Cys Gly Ser

35 40 45

Gly Ser Ser Gly Ser Ser

50

<210> 40

<211> 54

<212> PRT

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35 40 45

Gly Ser Ser Gly Ser Ser

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<210> 41

<211> 53

<212> PRT

<213> *Gymnea sylvestre*

<400> 41

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1 5 10 15

Val Lys Lys Asp Glu Leu Cys Met Trp Ser Arg Glu Val Cys Cys Glu

20 25 30

Pro Leu Glu Cys Tyr Tyr Thr Gly Trp Tyr Trp Ala Cys Gly Ser Gly

35 40 45

Ser Ser Gly Ser Ser

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<210> 42

<211> 54

<212> PRT

<213> Gymnea sylvestre

<400> 42

Met His His His His His His Ser Gly Ser Ser Ser Gly Ser Gly Cys

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Val Lys Lys Asp Glu Leu Cys Glu Leu Ala Val Asp Glu Cys Cys Glu

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30

Pro Leu Glu Cys Phe Gln Met Gly His Gly Phe Lys Arg Cys Gly Ser

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45

Gly Ser Ser Gly Ser Ser

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<210> 43

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